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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: Wed Aug 08 11:54:07 EDT 2007

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Application No: 10780638 Version No: 2.0

Input Set:

Output Set:

Started: 2007-08-07 17:13:50.435
Finished: 2007-08-07 17:13:52.249
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 814 ms
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Total Errors: 0
No. of SeqIDs Defined: 33
Actual SeqID Count: 33

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Error code	Error Description
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SEQUENCE LISTING

<110> Commonwealth Scientific and Industrial Research Organization
<120> Efficient gene silencing in plants using short dsRNA sequences
<130> BCS-03-2001
<140> 10780638
<141> 2004-02-19
<150> US 60/447,711
<151> 2003-02-19
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taatgggact caaaaataagg ctttggccca catcaaaaag ataagtaca aatcaaaaact 240
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ctctcttctt ctatctaaa tctgtgtata ctatccact attgggcttt ttattagtct 180
ataatgggac taaaaataag gctttggccc acataaaaaa gataagtac aaatcaaaac 240
taaattcaga gtctttctc ccacatcggt cactgtactc ttttgcgttt gtttatatat 300

tacacgaacc gatcttggt acgtcgagct aagtaacatg agcttgcatac ccatgtgggg 360
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<223> Pol III promoter region

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<223> poly T stretch

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<223> EcoRI restriction site

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agatctgttt tggctatgtt ggacgaaaca agtgaacttt taggatcaac tttagtttat 120
atatggagct tatatcgagc aataagataa gtggggcttt tatgttaattt aatgggctat 180
cgtccataga ttcactaata cccatgccca gtacccatgt atgcgtttca tataagctcc 240
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<223> XhoI restriction site

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<223> poly T stretch

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<222> (462)..(467)
<223> EcoRI restriction site

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atatggagct tatatcgagc aataagataa gtgggccttt tatgttaattt aatgggctat 120
cgtccataga ttcactaata cccatgccca gtaccatgt atgcgtttca tataagctcc 180
taatttctcc cacatcgctc aaatctaaac aaatcttgtt gtatatataa cactgaggga 240
gcaacattgg tcacgacctt acttgaacag gatctgtct ataggctcgt acctctgttt 300
ccttgatttc tcaagagaca ggcccttaac cctgggttgat gaaccatgac cgtgcggcta 360
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467

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cttgaaccgt agcttcgtt ttcttcttt taactttcca ttcggagttt ttgtatctg      180
tttcatagtt tgtccagga ttagaatgtt taggcatacga accttcaaga atttgatgt      240
ataaaaacatc ttcatctta agatatgtt ataattttca aaaggccccct gggaatctga      300
aagaagagaa gcaggcccat ttatatggaa aagaacaata gtatttctta tataggccca      360
tttaagttga aaacaatctt caaaaagtccc acatcgctta gataagaaaa cgaagctgag      420
tttatataca gctagagtcg actttttttt gagctc                                456

<210> 6
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<213> Artificial Sequence

<220>

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cttgaaccgt agcttcgtt ttcttcttt taacttcca ttccggagttt ttgtatctg 180

tttcatagtt tgtccagga ttagaatgat taggcatacga accttcaaga atttgattga 240

ataaaaacatc ttcatctta agatatgaag ataatttca aaaggcccct ggaaatctga 300

aagaagagaa gcagggccat ttatatggga aagaacaata gtatttctta tataggccca 360

tttaagttga aaacaatctt caaaagtccc acatcgctta gataagaaaa cgaagctgag 420

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<220>

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<220>

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<220>

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<222> (385)..(390)

<223> PvuI restriction site

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aggacaggcg tcttctactg gtgctaccag caaatgctgg aagccggaa cactgggtac 180

gttggaaacc acgtgatgtg gagtaagata aactgttagga gaaaagcatt tcgttagtgg 240

ccatgaagcc tttcaggaca tgtattgcag tatggccgg cccattacgc aattggacga 300

caacaaagac tagtattagt accacctcg ctatccacat agatcaaagc tggttaaaa 360

gagttgtgca gatgatccgt ggcacgatcg ttttttttg aattc 405

<210> 8

<211> 442

<212> DNA

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<220>

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gaattattat ggagttcatg gcaactgcta tggagtttt cctactggga attggaacgg      180
tttctacgaa attaactgtc cacacgttaa aaatataaat taatgcgtaa ttgttatttt      240
ttctataaca aataaaaaac tcaaatacga cataaatttt attacttaa ttgcacttta      300
gccttagaga tattgcgttg tagtcggcgt aggtgtgtca ggggccaata tattgttccc      360
acatcgccag tgcagcacat aaactctagc gttataagaa tctatccact atcaacggtc      420
acgatcggtt tttttgaat tc                                         442

<210>  9
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<223>  SalI restriction site

<220>

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gtggtgacgc atgtcgcgca agactgtaac cacgcgtctg ttgactggca ggtggtggcc      120
aatggtgatg tcagcggtga actgcgtgat gcggatcaac aggtggttgc aactggacaa      180
ggcactagcg ggatccagac gcgtggttac agtcttgcgc gacatgcgtc accacggtga      240
tatcgccac ccagggtgtc ggcgtggtgt agagcattac gctgcgatcg tcgac      295

<210> 10
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<220>
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<220>
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<223> GUS sequence (antisense)

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caccacgatg ccatgttcat ctgcccagtc gac                                93

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accgccacgg ccaccgccac taccccccac tccg 94

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<210> 16
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gccatatccc tacatggctg aaccacacca tcca 94

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<212> DNA
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<220>
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<220>
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atcatgtcac cacttccact agcagttgac aatg 94

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expression of the PHYB gene, derived from the 3' end of PHYB
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expression of the PHYB gene, derived from the 3' end of PHYB
(PHYB3hp21)-lower strand

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expression of the PDS gene (PDS42)-upper strand

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ccggtagga tcgtaatatt ccttacaagt taag 94

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